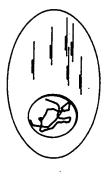


ERAFTSTO-≻ m

CORRESPONDING PROTEIN SYNTHESIS BLOCKED

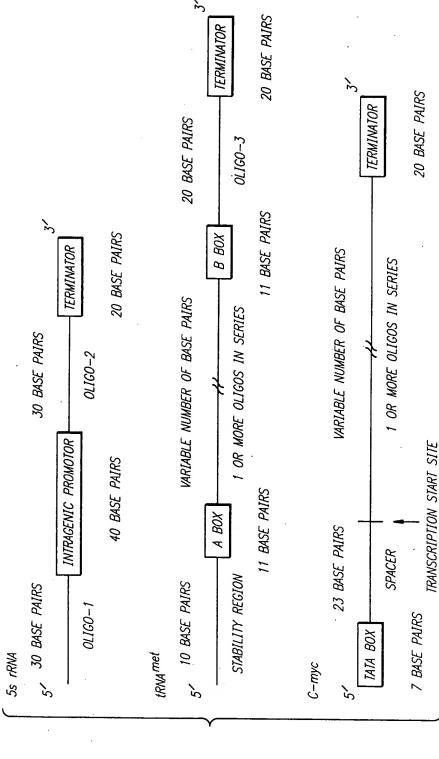


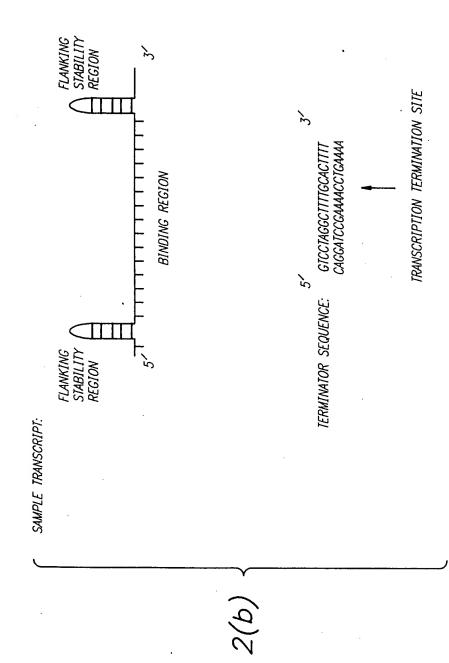
CHROMOSOMĘ OLIGONUCLEOTIDE RNA PROTEIN KEY: **-00**≺ TRANSCRIPTION UNIT RETROVIRAL VECTOR VIRAL PARTICLE

OLIGONUCLEOTIDE BINDS TO COMPLEMENTARY RNA

SPACETORS IN THE PLANTS







96.2 CRAFTCHALL <u>}:</u>

HERZ PROMOTER FRAGMENT

3' TCCTCTTCCTCCTCCTCCTCCTCCGACGAACTCCTTCATATTCTTA

- 5' AGGAGAAGGAGGAGGTGGAGGAGGGGCTGCTTGAGGAAGTATAAGAAT
- 5' UCCUCUUCCUCCUCCUCCUCCUCCC CU-RICH TRIPLEX FORMING RNA
- GA-RICH TRIPLEX FORMING RNA 3' AGGAGAAGGAGGAGGGGGAGGAGGG



FIG 4A

THE UB SMALL NUCLEAR RNA GENE

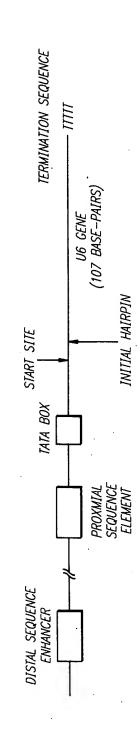


FIG. 4B

THE CHIMERIC OLIGONUCLEOTIDE PRODUCING GENE

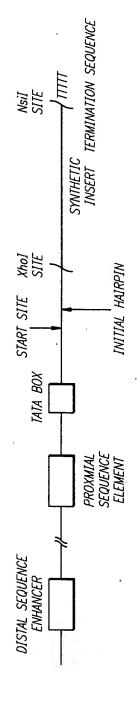
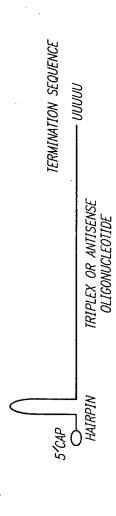
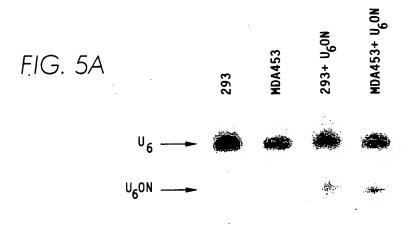
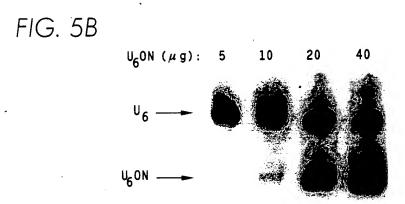
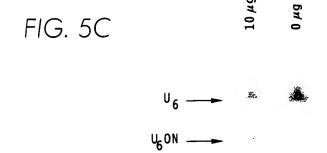


FIG. 4C









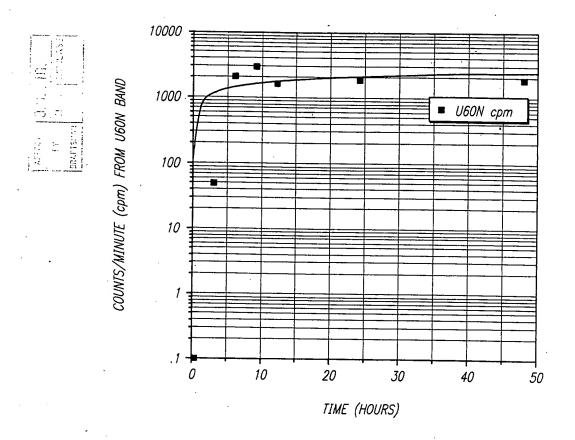


FIG. 6A

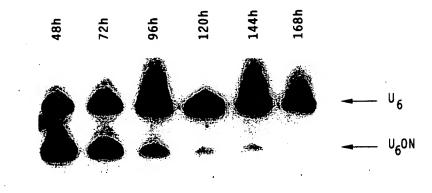


FIG. 6B

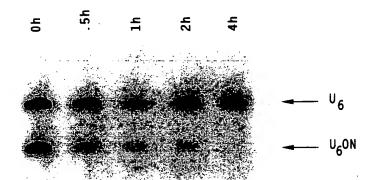


FIG. 7B

Ab: + + .

U₆ON: + .

FIG. 8

U₆ ---

U₆0N ____

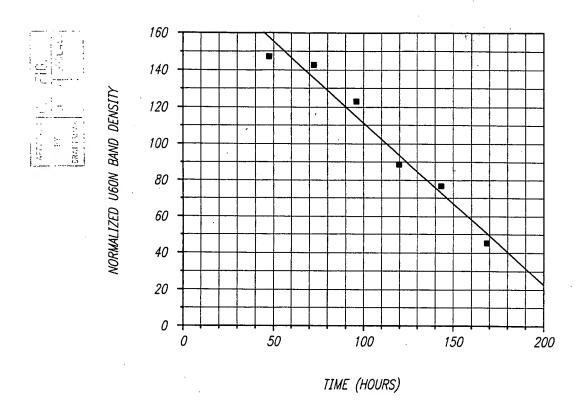
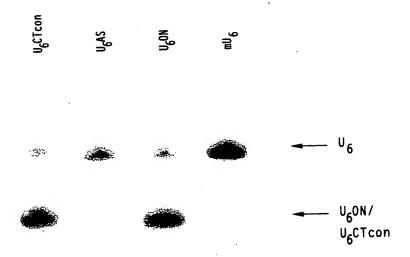


FIG. 7A

FIG. 9A

```
u c
  CG
c GC
  UA
  CG
  \mathsf{GC}
  UA20
{\tt 1} \ \mathsf{GCAUAUccu:} \mathsf{CGaccuccccuucccuucccuucccUUC:::C}
     UAUAccuuGC::::::::::::::::::::::::::::::GAAGuacGua
    U
                                                                60
    U 80
    U
                U6CTcon ENERGY = -12.72 kcal
                 (U60N ENERGY = -12.46 kcal)
1 20
GuGcuCGCUUCg:GCAgCACAUau:::CCuCGaC:::AUG<sup>a g</sup> c
C:CuuGCGAAGuaCGUaGUGUAagaacGG:GC:GgacUAC<sub>u u</sub>
 Α
                         60
 U
 A 80
 UUUUU
                 U6AS ENERGY = -30.83 kcal
                 (mU6 ENERGY = -26.48 kcal)
```



TRIPLEX RNA OLIGONUCLEOTIDE HER2 PROMOTER MAP. AND

TRIPLEX RNA OLIGONUCLEOTIDE

CCCAATCACAGGAGAAGGAGGAGGTGGAGGAGGAGGCTGCTTGAGGAAGTATAAGAA 3' GGGTTAGTGTCCTCCTCCTCCTCCTCCTCCTCCGACGACCTCCTTCATATTCTT e. nccncnnccnccncccccnccnccnccc 3.

CAAT

BOX

TATA ets-

ELEMENT BOX

FIG. 11A



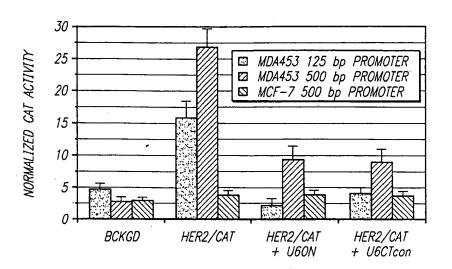


FIG. 11B

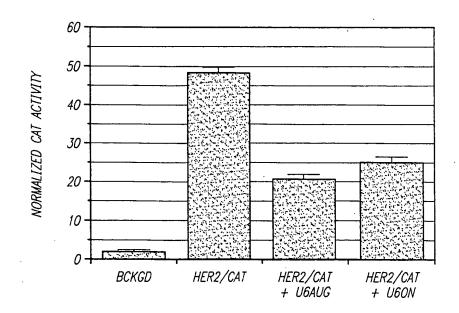
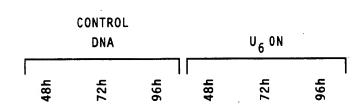
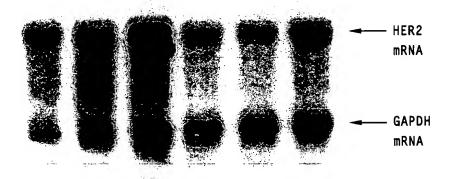


FIG. 12A





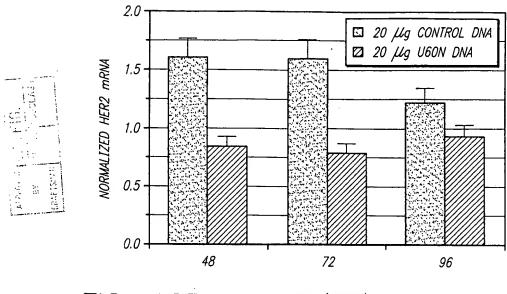


FIG. 12B

TIME (HOURS)

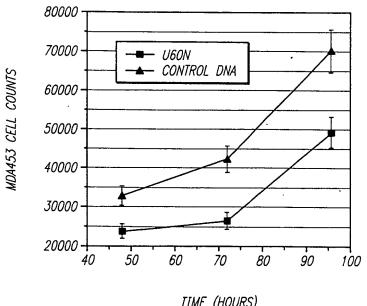


FIG. 12D

TIME (HOURS)

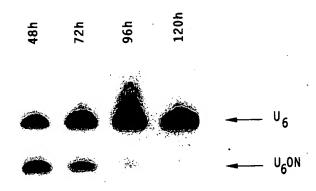


FIG. 12C

FIG. 14A

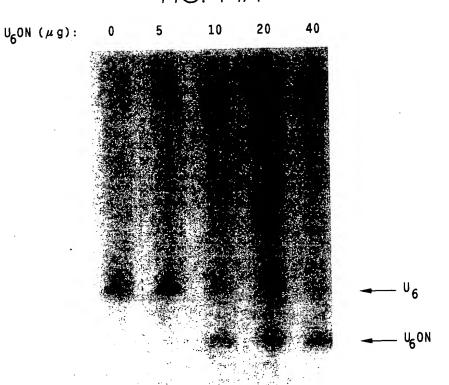
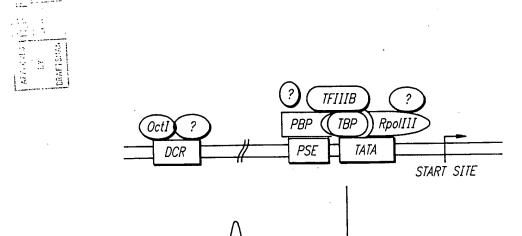


FIG. 13

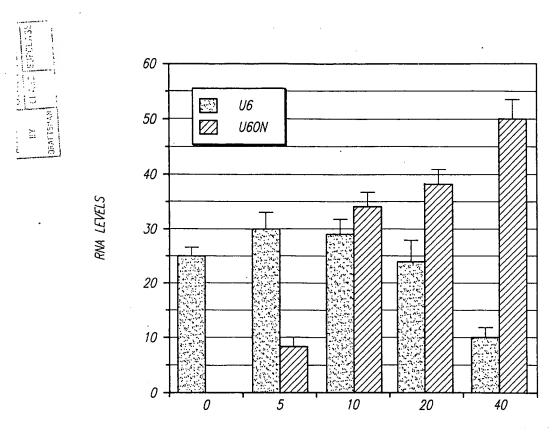


5' CAP

POSSIBLE FACTORS IN LIMITING SUPPLY:

RNA POLYMERASE III (RpolIII)
IFIIIB CONTAINING THE TATA BINDING PROTEIN (TBP)
PROXIMAL SEQUENCE ELEMENT BINDING PROTEIN (PBP)
UPSTREAM ENHANCERS (OctI,?)
OTHER UNCHARACTERIZED TRANSCRIPTIONAL FACTORS (?)
5' CAPPING ENZYME, CO-FACTORS
LUPUS ASSOCIATED ANTIGEN (La)

ON



TRANSFECTION DOSE (Lug)

FIG. 14B

FIG. 15

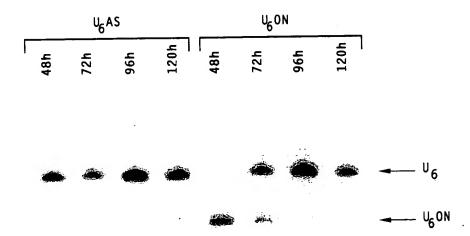
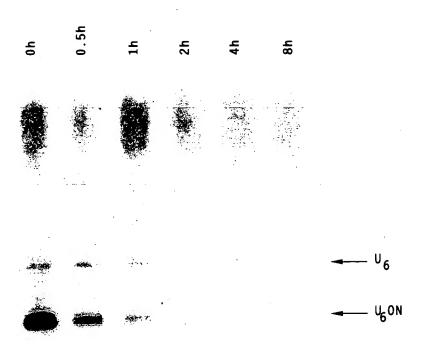


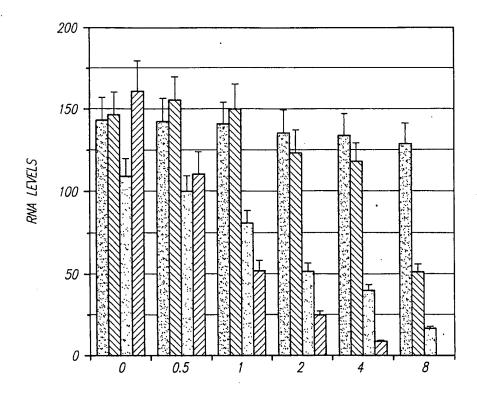
FIG. 16



MOST JULY, FIG.

FIG. 17

- 🔀 U6 (5μg U6ON)
- U6 (10µg U60N)
- U6 (20µg U60N)
- ∠ U60N (20µg U60N)



TIME (HOURS)

FIG. 18A

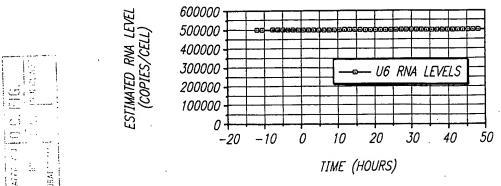


FIG. 18B

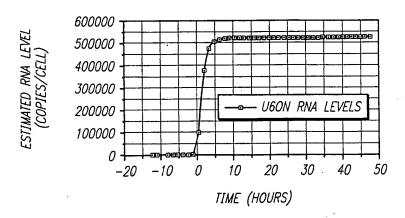


FIG. 18C

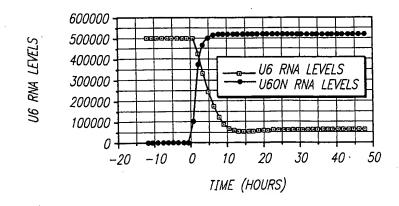


FIG. 19

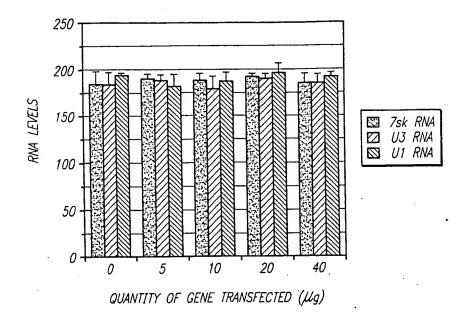


FIG. 20A

RNA LEVELS

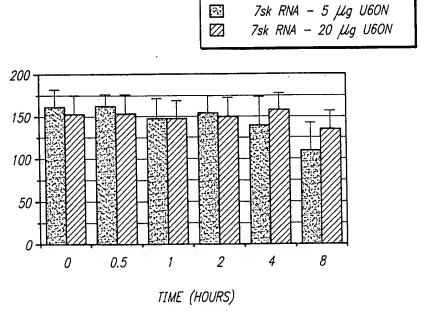


FIG. 20B

U1 RNA - 5 μg U60N
 U1 RNA - 20 μg U60N

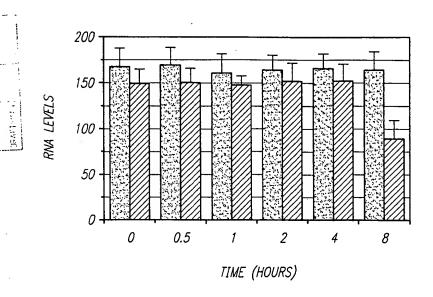
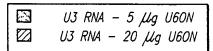
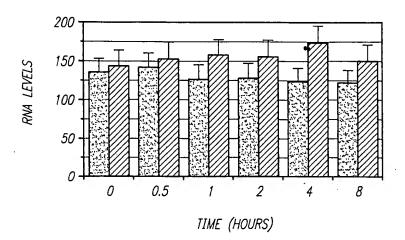
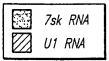


FIG. 20C







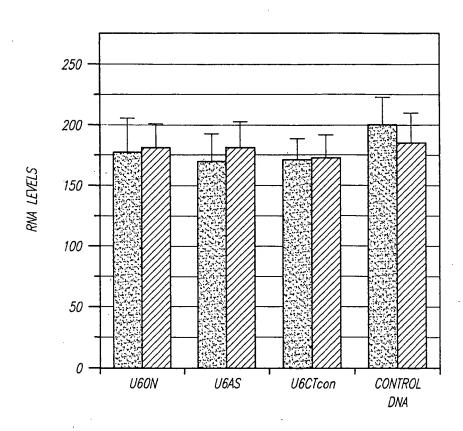


FIG. 21

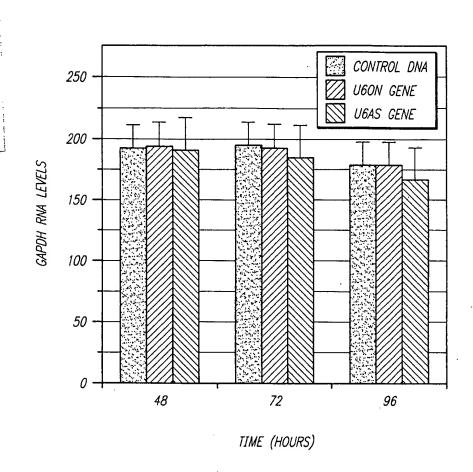


FIG. 22

FIG. 23A



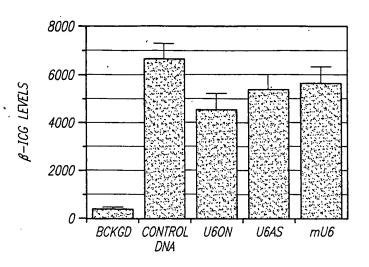
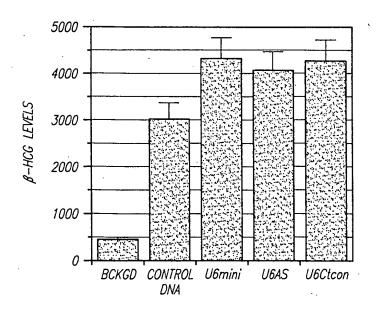


FIG. 23B



Asirana (1,17, F16, 34, F3)

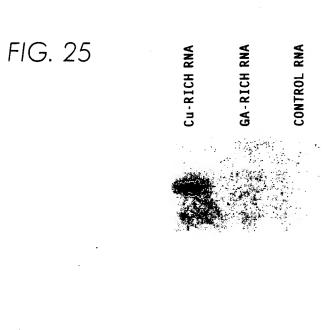
FIG. 24

3 GGTTAGTGTCCTCTTCCTCCTCCTCCTCCTCCTCCGACGAAC

5' CCAATCACAGGAGGAGGAGGAGGAGGAGGAGGAGGCTGCTTG

5' UCCUCUUCCUCCUCCCCCUCCUCCUCC... CU-RICH RNA

 5' GGGCCCCCCCCCGAGGUCGACGGUAUCG... CONTROL RNA



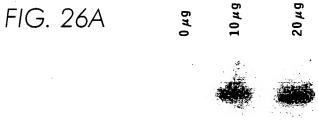


FIG. 26B

FIG. 27A

RY C.C.S. EM.CLASS.

U6 PARENT GENE

-240	TTCCCATGAT	TCCTTCATAT	TTGCATATAC
-210	GATACAAGGC	TGTTAGAGAG	ATAATTAGAA
-180	TTAATTTGAC	TGTAAACACA	AAGATATTAG
- 150	TACAAAATAC	GTGACGTAGA	AAGTAATAAT
-120	TTCTTGGGTA	GTTTGCAGTT	TTTAAAATTA
-90	TGTTTTAAAA	TGGACTATCA	TATGCTTACC
-60	GTAACTTGAA	AGTATTTCGA	TTTCTTGGCT
-30	TTATATATCT	TGTGGAAAGG	ACGAAACACC
+1	GTGCTCGCTT	CGGCAGCACA	TATCCTCGAG
+31	CATGGCCCCT	GCGCAAGGAT	GACACGCAAA
+61	TGCATGAAGC	GTTCCATATT	TTT 83 NUCLEOTIDES

FIG. 27B

U60N GENERATOR

-240	TTCCCATGAT	TCCTTCATAT	TTGCATATAC
-210	GATACAAGGC	TGTTAGAGAG	ATAATTAGAA
-180	TTAATTTGAC	TGTAAACACA	AAGATATTAG
-150	TACAAAATAC	GTGACGTAGA	AAGTAATAAT
-120	TTCTTGGGTA	GTTTGCAGTT	TTTAAAATTA
90	TGTTTTAAAA	TGGACTATCA	TATGCTTACC
-60	GTAACTTGAA	AGTATTTCGA	TTTCTTGGCT
-30	TTATATATCT	TGTGGAAAGG	ACGAAACACC
+1 .	GTGCTCGCTT	CGGCAGCACA	TATCCTCGAC
+31	TCCTCTTCCT	CCTCCACCTC	CTCCTCCCAT
+61	GCATGAAGCG	TTCCATATTT	TT 82 NUCLEOTIDES